

# SEQUENCE LISTING

<110> Cahoon, Rebecca E.  
Falco, Saverio Carl  
Pember, Stephen O.

<120> Chorismate Biosynthesis Enzymes

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<141> 2001-01-04

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Asp Ala Ile Asp Lys Val Arg Val Asn Gly Asn Ser Ile Gly Gly Val  
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Val Thr Cys Ile Ala Arg Asn Val Pro Arg Gly Leu Gly Ser Pro Val  
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Phe Asp Lys Leu Glu Ala Leu Leu Ala Lys Ala Met Leu Ser Leu Pro

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Tyr Gly Glu Ser His Gly Gly Gly Val Gly Cys Val Ile Ser Gly Cys
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Pro Pro Arg Ile Pro Leu Thr Glu Ala Asp Leu Gln Val Glu Leu Asp
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Arg Arg Arg Pro Gly Gln Ser Arg Ile Thr Ser Thr Arg Lys Glu Thr
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Asp Thr Cys Lys Ile Leu Ser Gly Thr His Glu Gly Val Thr Thr Gly
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Thr Pro Ile Leu Val Ile Val Pro Asn Thr Asp Gln Ile Gly Ser Asp
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Tyr Asp Phe Lys Tyr Gly Val Arg Ala Val Gln Gly Gly Gly Arg Ser
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Lys Val His Gln Val Val Leu Pro Glu Asp Ala Val Asp Tyr Gly Ser
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Val Thr Leu Glu Gln Ile Glu Ser Asn Ile Val Arg Cys Pro Asp Pro
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Glu Tyr Ala Glu Lys Met Ile Asp Ala Ile Asp Arg Val Arg Val Arg
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Gly Asp Ser Val Gly Gly Val Ile Thr Cys Val Ala Arg Asn Val Pro

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 Arg Pro Gly Gln Ser Arg Ile Thr Thr Pro Arg Lys Glu Thr Asp Thr  
                                  100                                 105                                 110  
 Cys Lys Ile Leu Ser Gly Thr His Glu Gly Met Thr Thr Gly Thr Pro  
                                  115                                 120                                 125  
 Ile His Val Phe Val Pro Asn Thr Asp Gln Arg Gly Gly Asp Tyr Ser  
           130                                 135                                 140  
 Glu Met Ala Lys Ala Tyr Arg Pro Ser His Ala Asp Ala Thr Tyr Asp  
          145                                 150                                 155                                 160  
 Phe Lys Tyr Gly Val Arg Ala Val Gln Gly Gly Gly Arg Ser Ser Ala  
                                  165                                 170                                 175  
 Arg Glu Thr Ile Gly Arg Val Ala Ala Gly Ala Leu Ala Lys Lys Ile  
                                  180                                 185                                 190  
 Leu Lys Leu Lys Ser Gly Val Glu Ile Leu Ala Phe Val Ser Lys Val  
          195                                 200                                 205  
 His Gln Val Val Leu Pro Glu Asp Ala Val Asp Tyr Asp Thr Val Thr  
          210                                 215                                 220  
 Met Glu Gln Ile Glu Ser Asn Ile Val Arg Cys Pro Asp Pro Glu Tyr  
          225                                 230                                 235                                 240  
 Ala Gln Lys Met Ile Asp Ala Leu Asp Lys Val Arg Val Arg Gly Asp  
                                  245                                 250                                 255  
 Ser Ile Gly Gly Val Val Thr Cys Ile Ala Arg Asn Val Pro Arg Gly  
                                  260                                 265                                 270  
 Ile Gly Ser Pro Val Phe Asp Lys Leu Glu Ala Glu Leu Ala Lys Ala  
          275                                 280                                 285  
 Met Leu Ser Leu Pro Ala Ser Lys Gly Phe Glu Ile Gly Ser Gly Phe  
          290                                 295                                 300

Val Phe Thr  
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cccgcctacc tccgactctc actcgcgtcct cgtcttccca agagacttca catacaggcg 180  
gctgggagta cctatggaaa tcactttcgt gttacaacat atggggaatc acatggagga 240  
ggtgttggtt gtgttattga tggatgtcct cctcgccttc ctctctctga agctgatatg 300  
caagtggatc ttgacagaag gaggccaggt cagagccgaa ttacaactcc tagaaaggag 360  
actgatacat gtaaaatatt ttcaggagtt tccgaaggaa tcactactgg nactccaatt 420  
catgtactgt acccanntac tgatcaanga gggcatgact atagcnagat ggnagtacnt 480  
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                   20                  25                  30  
 Leu Arg Leu Ser Leu Arg Pro Arg Leu Pro Lys Arg Leu His Ile Gln  
           35                  40                  45  
 Ala Ala Gly Ser Thr Tyr Gly Asn His Phe Arg Val Thr Thr Tyr Gly  
       50                  55                  60  
 Glu Ser His Gly Gly Gly Val Gly Cys Val Ile Asp Gly Cys Pro Pro  
   65                  70                  75                  80  
 Arg Leu Pro Leu Ser Glu Ala Asp Met Gln Val Asp Leu Asp Arg Arg  
                   85                  90                  95  
 Arg Pro Gly Gln Ser Arg Ile Thr Thr Pro Arg Lys Glu Thr Asp Thr  
                   100                  105                  110  
 Cys Lys Ile Phe Ser Gly Val Ser Glu Gly Ile Thr Thr Gly Thr Pro  
       115                  120                  125  
 Ile His Val Ser Val Pro Asn Thr Asp Gln Xaa Arg His Asp Tyr Ser  
       130                  135                  140  
 Glu Met Ala Leu Leu Ile Gly Leu His Ala Asn Ala Thr Tyr Asp Met  
   145                  150                  155                  160  
 Lys Tyr Gly Xaa Arg Ser Val Lys  
                   165